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Y^b ; the connecting group Y^b not being cleavable to release substrate R under conditions effective to cleave the second cleavage sites in the groups Y^1R and Y^2R and wherein:

- (i) the chemical fragment F^b contains a sensitising group G which sensitises the chemical fragment F^b to instrumental, e.g. mass spectroscopic analysis and/or:
- (ii) the fragment F^b contains a means for imparting a characteristic signature to the mass spectrum of the fragment.

13. (Amended) A chemical construct according to claim 1 wherein the sensitising group G is generated by cleavage at the first cleavage site of the group Y^1 or Y^2 or, when present, Y^a or the said cleavage site of Y^b .

14. (Amended) A chemical construct according to claim 1 wherein the sensitising group G is a basic amino group or a carboxylate group, preferably a basic amino group.

17. (Amended) A construct according to claim 1 wherein the fragment F^r and (where present) optionally the fragment F^a and (where present) optionally the fragment F^a contain a means for imparting a characteristic signature to the mass spectrum of the fragment.

20. (Amended) A chemical construct according to claim 18 wherein the isotopic label comprises an atom or atoms selected from $^1H/^2H$ (D), $^{79}Br/^{81}Br$, $^{12}C/^{13}C$, $^{14}N/^{15}N$ and $^{16}O/^{18}O$.

21. (Amended) A chemical construct according to claim 18 wherein the isotopic label(s) is/are located between the first and second cleavage sites of the groups Y^1 and Y^2 .

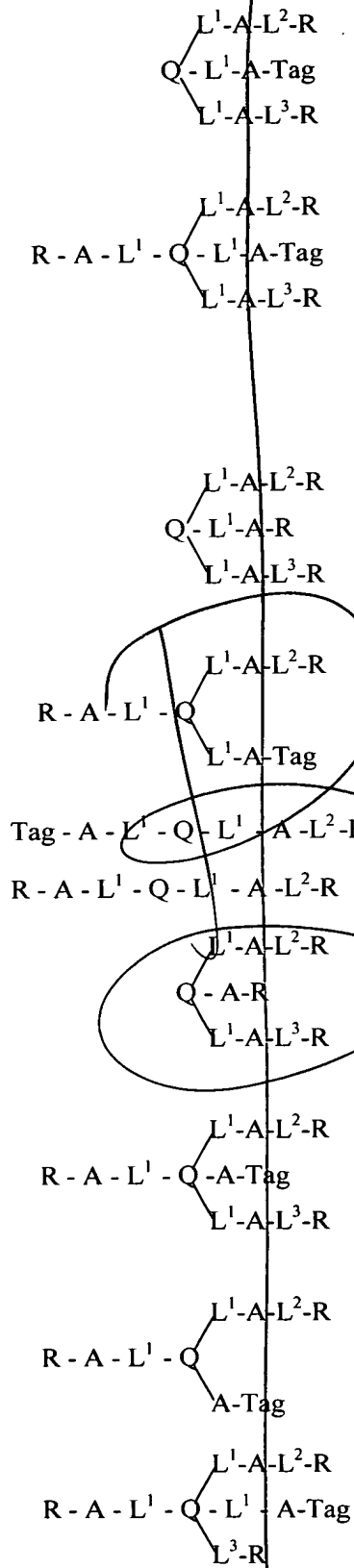
22. (Amended) A chemical construct as defined in claim 1 wherein the first and second cleavage sites in the groups Y^1 are defined by first and second linker groups L^1 and L^2 , first and second cleavage sites in the group Y^2 are defined by first and second linker groups L^1 and L^2 , the cleavage site in the group Y^a (where present) is defined by a linker group L^a and the cleavage site in the group Y^b (where present) is defined by a linker group L^b .

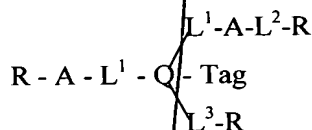
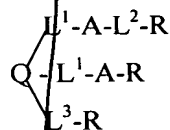
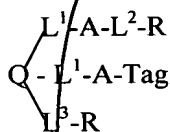
24. (Amended) A chemical construct according to claim 22 wherein a spacer group A is interposed between each pair of first and second linker groups, or between the linker group L^a and the coding tag, or between the linker group L^b and the substrate R, the spacer group A containing an isotopic peak splitting label.

25. (Amended) A chemical construct according to claim 1 having a formula selected from the group consisting of:

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wherein L^1 , L^2 , L^3 , A and R are as defined in claim 1 and "Tag" represents a coding sequence.

26. (Amended) A construct as claimed in claim 1 for use in a tiered release method of screening, the construct having the formula $Tag-A-L^1-Q-L^1-A-L^2-R$ wherein Tag, A, L^1 , Q, L^2 and R are as defined claim 1.
27. (Amended) A chemical construct according to claim 1 wherein the orthogonally cleavable cleavage sites can be cleaved by a reactions selected from acid catalysed cleavage, base catalysed cleavage, oxidative cleavage, reductive cleavage, nucleophilic displacement, electrophilic displacement, and thermal, photochemical and enzymatic cleavage.
28. (Amended) Intermediate chemical constructs for use preparing a chemical construct as defined in claim 1, the intermediate constructs having the formulae $Y^{1i}-Q-Y^{2i}$, $RY^{1i}-Q-Y^{2i}$ and $Y^{1i}-Q-Y^{2i}R$ wherein Y^{1i} and Y^{2i} are reactive or protected forms of the group Y; and R, Q and Y are as defined in claim 1.
29. (Amended) Intermediate constructs of the formulae $L^{2i}-A-L^1-Q-L^1-A^p$, $R-L^2-A-L^1-Q-L^1-A^p$, $L^{3i}-A-L^1-Q-L^1-A^p$, $R-L^3-A-L^1-Q-L^1-A^p$, $R-L^3-A-L^1-Q-L^1-A-L^{2i}$ and $L^{3i}-A-L^1-Q-L^1-A-L^2-R$ wherein L^{1i} , L^{2i} and L^{3i} are reactive or protected forms of the linker groups L^1 , L^2 and L^3 , A^p is a reactive or protected form of the spacer group A containing a peak splitting isotopic label, and Q, R, A, L^1 , L^2 and L^3 are as defined in claim 1.

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31. (Amended) An intermediate construct according to claim 29 wherein the solid support has bonded thereto a coding tag sequence L¹-A-Tag and/or a sequence R - A - L¹ -, or a precursor form thereof..

32. (Amended) A differential release method of assaying a chemical library for biological activity, the method comprising:

(i) subjecting a construct comprising a solid support Q having linked thereto groups Y¹R and Y²R as defined in claim 1 to cleavage conditions effective to release substrate R from the group Y¹R;

(ii) testing the substrate R released from the group Y¹R in a biological assay;

(iii) subsequently subjecting the construct to cleavage conditions effective to release substrate R from the group Y²R; and

(iv) testing the substrate R released from the group Y²R in a biological assay.

33. (Amended) A tiered release method of assaying a chemical library for biological activity, the method comprising:

(i) subjecting a construct as claimed in claim 1 to cleavage conditions effective to release a first portion of the substrate R from the group Y¹R;

(ii) testing the first portion of substrate R released from the group Y¹R in a biological assay;

(iii) subjecting the construct to cleavage conditions effective to release a second portion of the substrate R from the group Y¹R; and

(iv) testing the second portion of substrate R released from the group Y¹R in a biological assay.

34. A method of determining the identity of a substrate R linked to a solid support Q of a construct as claimed in claim 8 by mass spectrometric means; the solid support Q having a coding sequence attached thereto by means of a connecting group Y^a having a cleavage site cleavable to release a fragment F^a from the solid support, the fragment F^a comprising the coding sequence and at least a portion of the connecting group Y^a, wherein (i) the chemical fragment F^a contains a sensitising group G which sensitises the chemical fragment F^a to mass spectroscopic analysis;

the coding sequence comprising a sequence of coding groups the nature and order of which is indicative of the identity of the substrate R;

the method comprising cleaving the connecting group Y^a so as to release the fragment F^a from the solid support; subjecting the fragment Y^a to mass spectrometry under conditions effective to bring about mass spectral fragmentation of the coding group and the formation of mass spectral fragment ions corresponding to the loss of one

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